IN THE SPECIFICATION

Please substitute the following three paragraphs for those in the Brief Description of the Drawings section at pages 3-4, paragraphs 10-12.

- [10] Fig. 1A shows a schematic of assay for fractional allelic expression showing key steps. See text for additional details.
- [11] Fig. 2 <u>1B-1</u> and <u>1B-2</u> shows the result of allelic expression analyses performed as described below in note (3). Representative results are shown for eight genes. The shaded box represents approximated 95% confidence interval and red bars indicate individuals displaying significant variations, as defined in note (6).
- [12] Fig. 3 1C shows examples of two kindreds exhibiting Mendelian inheritance patterns in either the PKD2 or Calpain-10 gene. Only individuals who were heterozygous for the SNP or were used to deduce haplotypes are shown. The individuals displaying altered fractional allelic expression are shaded red, and the individuals originally found to display altered expression are indicated by arrows. An obligate carrier in the PKD2 pedigree who could not be scored is indicated with a red dot. The results of genotype analyses are shown directly above each member of the pedigrees. The markers employed are listed at the right and each allele observed in a family was assigned a number. Markers suggesting a recombination are underlined and the allele associated with altered expression is indicated in red. The fractional allelic expression data used to score the pedigree are shown above the genotype and were interpreted as described in the legend to Figure 2 IB-1 and IB-2.